## SEQUENCE LISTING

<110> Reed, John C.
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<120> Novel Card Domain Containing Polypeptides, Encoding Nucleic Acids, and Methods of Use

<130> P-LJ 4752

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tte cac etc etg gge cag eet etc tee cac ttg gee agg ege ett etg
Phe His Leu Leu Gly Gln Pro Leu Ser His Leu Ala Arg Arg Leu Leu
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Ser Leu Leu Arg Gly Arg Leu Pro Thr Leu Leu His Leu Gly Arg Leu
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Ala Leu Trp Gly Leu Gly Met Cys Cys Tyr Val Phe Ser Ala Gln Gln
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cgt g Arg G 45			-		_	_	-	-		-	-	_		_		194
gtt a Val I		_			_	_		-			_					242
tgt g Cys G			-					-		_	_					290
ccg a Pro I					tago	cttag	get t	agta	acaca	aa ga	actco	ccaat	t ta	ctatt	ttc	345
	ggca	ca g	atcaa	agctt	t go	ettt	ctaga	a aga	acgct	caa	agga	atato			aagccc aaagtt	
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aat agc cga gcc ctt att caa aga atg gga atg act gtt ata aag caa
Asn Ser Arg Ala Leu Ile Gln Arg Met Gly Met Thr Val Ile Lys Gln
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Ile Thr Asp Asp Leu Phe Val Trp Asn Val Leu Asn Arg Glu Glu Val
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		_				_	aca Thr		-		-	_	_	-	-	_	582
	-	_		_	-	_	tac Tyr						_				630
				-	-		gac Asp 125					_		_			678
					_		agg Arg	_	-					_			726
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		_		-			aaa Lys		-			_	_	-	-		822
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÷							ctc Leu 205										918
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475

480

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		-												gcc Ala			1926
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Left from Green world bank														att Ile			2118
	_													gca Ala			2166
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Cys Lys Asn Ile Tyr Ser Leu Met Val Glu Ala Ser Pro Leu Thr Ile 715 720
gaa gat gag agg cac atc aca tct gta aca aac ctg aaa acc ttg agt 2502  Glu Asp Glu Arg His Ile Thr Ser Val Thr Asn Leu Lys Thr Leu Ser  730  735  740
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Met Asn Glu Glu Asp Ala Ile Lys Leu Ala Glu Gly Leu Lys Asn Leu  780  785
795 Robert His Leu Ser Asp Ile Gly Glu
Gly Met Asp Tyr Ile Val Lys Ser Leu Ser Ser Glu Pro Cys Asp Leu 810 815
02= 4 1
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Leu Ile Asp Arg Met Asn Val Leu Glu Gln Leu Thr Ala Leu Met Leu  875  880
Pro Trp Gly Cys Asp Val Gln Gly Ser Leu Ser Ser Leu Leu Lys His
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cct ctg aaa Pro Leu Lys 935		_										3126
agt gat gga Ser Asp Gly		Ala P										3174
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att aca ggt Ile Thr Gly 1015						taaa	ataaa	agt (	gtaci	cgaa	ag	3368
Ile Thr Gly	Ala Phe	Lys L 1020	eu Val			taaa	ataaa	agt g	gtaci	cgaa	ag	3368 3396
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Ala	Phe	Phe		Gly	Lys	Ser	Leu		Ile	Asn	Ser	Gly		Ile	Pro
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Gln	Glu	Phe	Ara	645 Thr	Leu	Glu	Val	Thr	650 Leu	Arg	Asp	Phe	Ser	655 Lys	Leu
			660					665					670		
Asn	Lys	675	Asp	116	Arg	Tyr	680	GIY	гÀг	11e	Pne	685	ser	Ата	THE
Ser	Leu 690	Arg	Leu	Gln	Ile	Lys 695	Arg	Суѕ	Ala	Gly	Val 700	Ala	Gly	Ser	Leu
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Glv	Glv	T.em	740 Thr	Asn	Ser	Len	Glv	745 Asn	Leu	Lvs	Asn	Len	750 Thr	Lvs	Leu
•	•	755		-			760			-		765		_	
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Ser	Glu	Pro			Leu	Glu	Glu			Leu	Val	Ser			Leu
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Lvs	Leu	835 Ser	Ile	Leu	Asp	Leu	840 Ser	Glu	Asn	Tvr	Leu	845 Glu	Lvs	Asp	Glv
-	850				_	855				_	860				_
865					870					875					880
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945	-		_		950		-	-	_	955				-	960
	Ser Asn Ser Leu Ser Leu Ser Leu Ser Can Asn Ser Can Asn Can Can Can Can Can Can Can Can Can Ca	545         Leu           Ala         Phe           Asp         Tyr           Leu         Asp           Glu         Lys           625         Thr           Glu         Thr           Glu         Cys           Ser         Leu           690         Ser           Asn         Leu           Gly         Gly           Ile         Met           770         Glu           Glu         Ser           Leu         770           Glu         Ser           Leu         770           Glu         Ser           Jeu         Ser           Leu         770           Glu         Ser           Jeu         Ser           Jeu         Ser           Jeu         Ser           Jeu         Ser           Jeu         Thr           Ser         Jeu           Jeu         Thr           Ser         Jeu           Jeu         Jeu           Jeu         Jeu           Jeu         Jeu           Jeu	11e         Leu         Lys           545         Tyr         Gln           Ala         Phe         Phe           Asp         Phe         Phe           Glu         Lys         Ala           G25         Thr         Tyr           Glu         Lys         Ala           G25         Thr         Tyr           Glu         Clu         Phe           Asn         Lys         Gln           Asp         675         Arg           690         Arg         Arg           690         Pro         Arg           Asn         Leu         755           Ala         Ser         Pro           Asn         Leu         755           Leu         Asp         85           Leu         Asp         85 </td <td>11e         Leu         Lys         Ala           545         Tyr         Gln         Glu           Leu         Phe         580           Asp         Tyr         Leu         Phe           580         Asp         Phe         580           Asp         Tyr         Leu         Phe           610         Leu         Ala         Ala           610         Tyr         Ile           610         Tyr         Ile           610         Tyr         Ile           610         Tyr         Ile           620         Arg         Leu           Asn         Lys         Arg           Asn         Leu         Thr           705         Ala         Ser           Ala         Ser         Pro         Leu           770         Leu         Lys           11e         Met         Asp         Asn           12e         Asp         Ile           Ser         Ala         Asn         Ala           12e         Asp         Ile           12e         Asp         Ile           12e         Asp</td> <td>11e         Leu         Lys         Ala         11e           545         12e         565           Leu         Tyr         Gln         Gln         Gly           Asp         Tyr         Leu         Phe         Asp           Leu         Asp         Phe         Ile         Lys           Glu         Lys         Ala         Glu         Glu           Glu         Lys         Ala         Glu       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  Lease   Le	The color of the	The color of the	The Leu   Leu   Sys   Ala   She   She	The Leu Lys	The Leu Lys	The Leu Lys	The Leu

Phe Glu Asn Leu Lys Gln Leu Val Phe Phe Asp Phe Ser Thr Lys Glu 970 965 Phe Leu Pro Asp Pro Ala Leu Val Arg Lys Leu Ser Gln Val Leu Ser 985 Lys Leu Thr Phe Leu Gln Glu Ala Arg Leu Val Gly Trp Gln Phe Asp 1000 Asp Asp Asp Leu Ser Val Ile Thr Gly Ala Phe Lys Leu Val Thr Ala 1010 1015 <210> 98 <211> 1395 <212> DNA <213> Homo sapien <220> <221> CDS <222> (277) . . . (1353) <400> 98 cgcccgggca ggtgtttata ctccggaggg tgtccccgtg cgtcatcggt ggagtggacc 60 aaaactggtg atctgtttgc cctgtgtgac cttgcccaga accctgctga ctgagagaac 120 acatetgetg gaagteetet gggatteaag gtacagggaa tgaagagtag tittacagaa 180 aaaagaggac aatattggga tcacctttga cctttccatt tggaaataat attttctatt 240 gtgttataga aaggtgggaa gctttcatcc agaaca atg aat ttc ata aag gac Met Asn Phe Ile Lys Asp aat agc cga gcc ctt att caa aga atg gga atg act gtt ata aag caa 342 Asn Ser Arg Ala Leu Ile Gln Arg Met Gly Met Thr Val Ile Lys Gln atc aca gat gac cta ttt gta tgg aat gtt ctg aat cgc gaa gaa gta 390 Ile Thr Asp Asp Leu Phe Val Trp Asn Val Leu Asn Arg Glu Glu Val 25 30 aac atc att tgc tgc gag aag gtg gag cag gat gct gct aga ggg atc 438 Asn Ile Ile Cys Cys Glu Lys Val Glu Gln Asp Ala Ala Arg Gly Ile 40 att cac atg att ttg aaa aag ggt tca gag tcc tgt aac ctc ttt ctt 486 Ile His Met Ile Leu Lys Lys Gly Ser Glu Ser Cys Asn Leu Phe Leu 55 65 60 aaa too ott aag gag tgg aac tat oot ota ttt cag gac ttg aat gga 534 Lys Ser Leu Lys Glu Trp Asn Tyr Pro Leu Phe Gln Asp Leu Asn Gly caa agt ggt ctg act gac agc ttg ggt aac ttg aag aac ctt aca aag 582 Gln Ser Gly Leu Thr Asp Ser Leu Gly Asn Leu Lys Asn Leu Thr Lys 100 ctc ata atg gat aac ata aag atg aat gaa gaa gat gct ata aaa cta

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										gat Asp							726
					-				_	att Ile 160			-		_	-	774
										gct Ala	_					_	822
	-		•	_			_			gaa Glu			_	_			870
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<b>±</b>			_				_			gat Asp				_			1062
		_							-	aaa Lys			_	_	_		1110
										gga Gly							1158
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	_				_		_		_	aga Arg			_				1254

325

315

ā T

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8

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Val Pro Gln Leu Val Lys Leu Gly Leu Lys Asn Trp Arg Leu Thr Asp

215

230

245

220

235

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Ser Pro Lys Thr Thr Thr Asp Met Tyr Leu Leu Ile Leu Gln His Phe 485 490 Leu Leu His Ala Thr Pro Pro Asp Ser Ala Ser Gln Gly Leu Gly Pro 505 Ser Leu Leu Arg Gly Arg Leu Pro Thr Leu Leu His Leu Gly Arg Leu 520 Ala Leu Trp Gly Leu Gly Met Cys Cys Tyr Val Phe Ser Ala Gln Gln 535 Leu Gln Ala Ala Gln Val Ser Pro Asp Asp Ile Ser Leu Gly Phe Leu 550 Val Arg Ala Lys Gly Val Val Pro Gly Ser Thr Ala Pro Leu Glu Phe 565 570 · Leu His Ile Thr Phe Gln Cys Phe Phe Ala Ala Phe Tyr Leu Ala Leu 585 Ser Ala Asp Val Pro Pro Ala Leu Leu Arg His Leu Phe Asn Cys Gly 600 Arg Pro Gly Asn Ser Pro Met Ala Arg Leu Leu Pro Thr Met Cys Ile 615 Gln Ala Ser Glu Gly Lys Asp Ser Ser Val Ala Ala Leu Leu Gln Lys 620 630 635 Ala Glu Pro His Asn Leu Gln Ile Thr Ala Ala Phe Leu Ala Gly Leu 650 Leu Ser Arg Glu His Trp Gly Leu Leu Ala Glu Cys Gln Thr Ser Glu 665 Lys Ala Leu Leu Arg Arg Gln Ala Cys Ala Arg Trp Cys Leu Ala Arg 680 Ser Leu Arg Lys His Phe His Ser Ile Pro Pro Ala Ala Pro Gly Glu 695 Ala Lys Ser Val His Ala Met Pro Gly Phe Ile Trp Leu Ile Arg Ser 710 Leu Tyr Glu Met Gln Glu Glu Arg Leu Ala Arg Lys Ala Ala Arg Gly 725 730 Leu Asn Val Gly His Leu Lys Leu Thr Phe Cys Ser Val Gly Pro Thr 740 745 Glu Cys Ala Ala Leu Ala Phe Val Leu Gln His Leu Arg Arg Pro Val 760 Ala Leu Gln Leu Asp Tyr Asn Ser Val Gly Asp Ile Gly Val Glu Gln 775 Leu Leu Pro Cys Leu Gly Val Cys Lys Ala Leu Tyr Leu Arg Asp Asn 790 Asn Ile Ser Asp Arg Gly Ile Cys Lys Leu Ile Glu Cys Ala Leu His 805 810 Cys Glu Gln Leu Gln Lys Leu Ala Leu Gly Asn Asn Tyr Ile Thr Ala 825 Ala Gly Ala Gln Val Leu Ala Glu Gly Leu Arg Gly Asn Thr Ser Leu 840 Gln Phe Leu Gly Phe Trp Gly Asn Arg Val Gly Asp Glu Gly Ala Gln 855 Ala Leu Ala Glu Ala Leu Gly Asp His Gln Ser Leu Arg Trp Leu Ser 870 875 Leu Val Gly Asn Asn Ile Gly Ser Val Gly Ala Gln Ala Leu Ala Leu 890 Met Leu Ala Lys Asn Val Met Leu Glu Glu Leu Cys Leu Glu Glu Asn

905

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His Leu Gln Asp Glu Gly Val Cys Ser Leu Ala Glu Gly Leu Lys Lys
          915 920 925
    Asn Ser Ser Leu Lys Ile Leu Asn Ile Lys Ile His Ala Ser Gly Phe
       930 935
                             940
    Asn Lys Leu Leu Glu Ser Ile Phe Cys Ile Leu Leu Val Val Glu Ala
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n	<223> primer	
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=±	210. 140	
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Glu Ile Leu Gln His Asp Pro Asp Ser Ile Leu Asp Thr Leu Thr Ser
                                25
Arg Arg Leu Ile Ser Glu Glu Glu Tyr Glu Thr Leu Glu Asn Val Thr
Asp Leu Leu Lys Lys Ser Arg Lys Leu Leu Ile Leu Val Gln Lys Lys
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Gly Glu Ala Thr Cys Gln His Phe Leu Lys Cys
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Met Cys Ser Gln Glu Ala Phe Gln Ala Gln Arg Ser Gln Leu Val Glu
1
ctg ctg gtc tca ggg tcc ctg gaa ggc ttc gag agt gtc ctg gac tgg
                                                                   96
Leu Leu Val Ser Gly Ser Leu Glu Gly Phe Glu Ser Val Leu Asp Trp
             20
                                 25
ctg ctg tcc tgg gag gtc ctc tcc tgg gag gac tac gag ggc ttc cac
                                                                   144
Leu Leu Ser Trp Glu Val Leu Ser Trp Glu Asp Tyr Glu Gly Phe His
ctc ctg ggc cag cct ctc tcc cac ttg gcc agg cgc ctt ctg gac acc
                                                                   192
Leu Leu Gly Gln Pro Leu Ser His Leu Ala Arg Arg Leu Leu Asp Thr
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75

228

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Val Trp Asn Lys Gly Thr Trp Ala Cys Gln Lys Leu 70

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Leu Leu Ser Trp Glu Val Leu Ser Trp Glu Asp Tyr Glu Gly Phe His
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Leu Leu Gly Gln Pro Leu Ser His Leu Ala Arg Arg Leu Leu Asp Thr
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Val Trp Asn Lys Gly Thr Trp Ala Cys Gln Lys Leu
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Pro Ala Arg Asp Leu Gln Ser His Arg Pro Ala Ile Val Arg Arg Leu
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                 5
                                                         15
cac age cat gtg gag aac atg ctg gac ctg gca tgg gag cgg ggt ttc
                                                                   96
His Ser His Val Glu Asn Met Leu Asp Leu Ala Trp Glu Arg Gly Phe
             20
gtc agc cag tat gaa tgt gat gaa atc agg ttg ccg atc ttc aca ccg
                                                                   144
Val Ser Gln Tyr Glu Cys Asp Glu Ile Arg Leu Pro Ile Phe Thr Pro
         35
                             40
tee cag agg gea aga agg etg ett gat ett gee aeg gtg aaa geg aat
                                                                   192
Ser Gln Arg Ala Arg Arg Leu Leu Asp Leu Ala Thr Val Lys Ala Asn
                         55
gga ttg gct gcc ttc ctt cta caa cat gtt cag gaa tta cca gtc cca
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Gly Leu Ala Ala Phe Leu Leu Gln His Val Gln Glu Leu Pro Val Pro
65
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ttq
                                                                   243
Leu
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65

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<211> 888 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (1) ... (888) gac gat gcg gac act gtg ctg gtg gtg ggt gag gcg ggc agt ggc aag 48 Asp Asp Ala Asp Thr Val Leu Val Val Gly Glu Ala Gly Ser Gly Lys 15 1 age acg etc etg cag egg etg cae ttg etg tgg get gea ggg caa gae 96 Ser Thr Leu Leu Gln Arg Leu His Leu Leu Trp Ala Ala Gly Gln Asp 20 tto cag gaa ttt ctc ttt gtc ttc cca ttc agc tgc cgg cag ctg cag 144 Phe Gln Glu Phe Leu Phe Val Phe Pro Phe Ser Cys Arg Gln Leu Gln 40 tgc atg gcc aaa cca ctc tct gtg cgg act cta ctc ttt gag cac tgc 192 Cys Met Ala Lys Pro Leu Ser Val Arg Thr Leu Leu Phe Glu His Cys 55 tgt tgg cct gat gtt ggt caa gaa gac atc ttc cag tta ctc ctt gac 240 Cys Trp Pro Asp Val Gly Gln Glu Asp Ile Phe Gln Leu Leu Asp

cac cct gac cgt gtc ctg tta acc ttt gat ggc ttt gac gag ttc aag

His Pro Asp Arg Val Leu Leu Thr Phe Asp Gly Phe Asp Glu Phe Lys

90

288

95

70

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		_	_		_					_	_			_	ctg Leu	_	384
		-		_		-					-	-	-		gcg Ala		432
			_			_						_			tct Ser	_	480
	_				_		_		-	_					999 Gly 175		528
that that the		_	_			_	_						-	_	cac His		576
The first of the	_	_		_		_				_				_	cac His	_	624
Took lieur															gat Asp		672
ii" Saali Qenn Unn		_	_		_	_			_	_		-			cca Pro	_	720
		_			-	_			_					_	ctc Leu 255		768
					_	_									atg Met		816
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	-	_					ttc Phe 295										888

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Cys Met Ala Lys Pro Leu Ser Val Arg Thr Leu Leu Phe Glu His Cys
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Cys Trp Pro Asp Val Gly Gln Glu Asp Ile Phe Gln Leu Leu Leu Asp
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                    70
His Pro Asp Arg Val Leu Leu Thr Phe Asp Gly Phe Asp Glu Phe Lys
Phe Arg Phe Thr Asp Arg Glu Arg His Cys Ser Pro Thr Asp Pro Thr
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Ser Val Gln Thr Leu Leu Phe Asn Leu Leu Gln Gly Asn Leu Leu Lys
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Asn Ala Arg Lys Val Val Thr Ser Arg Pro Ala Ala Val Ser Ala Phe
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Leu Arg Lys Tyr Ile Arg Thr Glu Phe Asn Leu Lys Gly Phe Ser Glu
                                       155
                    150
Gln Gly Ile Glu Leu Tyr Leu Arg Lys Arg His His Glu Pro Gly Val
                                    170
Ala Asp Arg Leu Ile Arg Leu Leu Gln Glu Thr Ser Ala Leu His Gly
                                185
Leu Cys His Leu Pro Val Phe Ser Trp Met Val Ser Lys Cys His Gln
                            200
Glu Leu Leu Gln Glu Gly Gly Ser Pro Lys Thr Thr Asp Met
                                            220
                        215
Tyr Leu Leu Ile Leu Gln His Phe Leu Leu His Ala Thr Pro Pro Asp
                                        235
Ser Ala Ser Gln Gly Leu Gly Pro Ser Leu Leu Arg Gly Arg Leu Pro
                                    250
Thr Leu Leu His Leu Gly Arg Leu Ala Leu Trp Gly Leu Gly Met Cys
                                265
Cys Tyr Val Phe Ser Ala Gln Gln Leu Gln Ala Ala Gln Val Ser Pro
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                            280
Asp Asp Ile Ser Leu Gly Phe Leu
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														cat His		192
														ctt Leu		240
														tgg Trp 95		288
														gca Ala		336
														gcc Ala		384
ggt Gly	gtc Val 130	gtg Val	cca Pro	Gly 333	agt Ser	acg Thr 135	gcg Ala	ccc Pro	ctg Leu	gaa Glu	ttc Phe 140	ctt Leu	cac His	atc Ile	act Thr	432
														gat Asp		480
										Cys				ggc Gly 175		528
tca Ser	cca Pro	atg Met	gcc Ala 180	Arg	ctc Leu	ctg Leu	ccc Pro	acg Thr 185	atg Met	tgc Cys	atc Ile	cag Gln	gcc Ala 190	tcg Ser	gag Glu	576
gga Gly	aag Lys	gac Asp	agc Ser	agc Ser	gtg Val	gca Ala	gct Ala	ttg . Leu	ctg Leu	cag Gln	aag Lys	gcc Ala	gag Glu	ccg Pro	cac His	624

1209

200

195

gtg ctg gcc

Val Leu Ala

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340 345 350 Leu Gly Val Cys Lys Ala Leu Tyr Leu Arg Asp Asn Asn Ile Ser Asp

360 355 Arg Gly Ile Cys Lys Leu Ile Glu Cys Ala Leu His Cys Glu Gln Leu 375 Gln Lys Leu Ala Leu Gly Asn Asn Tyr Ile Thr Ala Ala Gly Ala Gln 395 390 Val Leu Ala <210> 177 <211> 261 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (1) ... (261) <400> 177 atg aat ttc ata aag gac aat agc cga gcc ctt att caa aga atg gga Met Asn Phe Ile Lys Asp Asn Ser Arg Ala Leu Ile Gln Arg Met Gly 10 5 atg act gtt ata aag caa atc aca gat gac cta ttt gta tgg aat gtt 96 Met Thr Val Ile Lys Gln Ile Thr Asp Asp Leu Phe Val Trp Asn Val 25 20 ctg aat cgc gaa gaa gta aac atc att tgc tgc gag aag gtg gag cag Leu Asn Arg Glu Glu Val Asn Ile Ile Cys Cys Glu Lys Val Glu Gln 40 35 gat gct gct aga ggg atc att cac atg att ttg aaa aag ggt tca gag 192 Asp Ala Ala Arg Gly Ile Ile His Met Ile Leu Lys Lys Gly Ser Glu 55 tcc tgt aac ctc ttt ctt aaa tcc ctt aag gag tgg aac tat cct cta 240 Ser Cys Asn Leu Phe Leu Lys Ser Leu Lys Glu Trp Asn Tyr Pro Leu 261 ttt cag gac ttg aat gga caa Phe Gln Asp Leu Asn Gly Gln 85 <210> 178 <211> 87 <212> PRT <213> Homo sapiens <400> 178 Met Asn Phe Ile Lys Asp Asn Ser Arg Ala Leu Ile Gln Arg Met Gly 10 Met Thr Val Ile Lys Gln Ile Thr Asp Asp Leu Phe Val Trp Asn Val

120

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cag ttt ggt Gln Phe Gly 130							432
gcc cag gct Ala Gln Ala 145	Leu Ile A		_		_	-	480
ttg ttg ctc Leu Leu Leu		•					528
acc cct ctc Thr Pro Leu			_				576
gag ttc cac Glu Phe His 195			~ ~			•	624
ctg ttg ata Leu Leu Ile 210	_					_	672
gac ttc att Asp Phe Ile 225	Arg Ser L						720
gtg ttc tcc Val Phe Ser	•	-	_				768
aat gag gat Asn Glu Asp	_	_		_			816
caa agg ttc Gln Arg Phe 275							864
tac aca gca Tyr Thr Ala 290							891
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Ala Leu Thr Lys Phe Lys Phe Val Phe Phe Leu Arg Leu Ser Arg Ala
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Gln Gly Gly Leu Phe Glu Thr Leu Cys Asp Gln Leu Leu Asp Ile Pro
Gly Thr Ile Arg Lys Gln Thr Phe Met Ala Met Leu Leu Lys Leu Arg
                                        75
                    70
Gln Arg Val Leu Phe Leu Leu Asp Gly Tyr Asn Glu Phe Lys Pro Gln
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Asn Cys Pro Glu Ile Glu Ala Leu Ile Lys Glu Asn His Arg Phe Lys
                               105
Asn Met Val Ile Val Thr Thr Thr Glu Cys Leu Arg His Ile Arg
                            120
Gln Phe Gly Ala Leu Thr Ala Glu Val Gly Asp Met Thr Glu Asp Ser
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                       135
Ala Gln Ala Leu Ile Arg Glu Val Leu Ile Lys Glu Leu Ala Glu Gly
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Leu Leu Gln Ile Gln Lys Ser Arg Cys Leu Arg Asn Leu Met Lys
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Thr Pro Leu Phe Val Val Ile Thr Cys Ala Ile Gln Met Gly Glu Ser
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                                                    190
           180
Glu Phe His Ser His Thr Gln Thr Thr Leu Phe His Thr Phe Tyr Asp
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Leu Leu Ile Gln Lys Asn Lys His Lys His Lys Gly Val Ala Ala Ser
                        215
                                            220
Asp Phe Ile Arg Ser Leu Asp His Arg Gly Asp Leu Ala Leu Glu Gly
                   230
                                        235
Val Phe Ser His Lys Phe Asp Phe Glu Leu Gln Asp Val Ser Ser Val
                                    250
Asn Glu Asp Val Leu Leu Thr Thr Gly Leu Leu Cys Lys Tyr Thr Ala
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Gln Arg Phe Lys Pro Lys Tyr Lys Phe Phe His Lys Ser Phe Gln Glu
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Tyr Thr Ala Gly Arg Arg Leu Ser Ser
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Gly Asn Leu Lys Asn Leu Thr Lys Leu Ile Met Asp Asn Ile Lys Met
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aat qaa qaa qat qct ata aaa cta gct gaa ggc ctg aaa aac ctg aag

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	_	_	_				_			_		gac Asp					144
	_	_			-	_		_		-	_	ccc Pro 60	_	_		_	192
												aat Asn					240
		_	-					-	-		-	agc Ser			_		288
and Said South And		_			_	_		_			~	gct Ala			_		336
4 min a m 7 min 1				_				_	_			gca Ala	-	-			384
17 July 18 18 18 18 18 18 18 18 18 18 18 18 18												ctg Leu 140	_			_	432
dan tasi tasi												aaa Lys			_		480
:										-		ttt Phe		_			528
												aat Asn				-	576
								_	_			aat Asn		_			618

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<212> PRT

<213> Homo sapiens

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Lys Met Cys Leu Phe His Leu Thr His Leu Ser Asp Ile Gly Glu Gly
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Met Asp Tyr Ile Val Lys Ser Leu Ser Ser Glu Pro Cys Asp Leu Glu
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Glu Ile Gln Leu Val Ser Cys Cys Leu Ser Ala Asn Ala Val Lys Ile
Leu Ala Gln Asn Leu His Asn Leu Val Lys Leu Ser Ile Leu Asp Leu
                                   90
               85
Ser Glu Asn Tyr Leu Glu Lys Asp Gly Asn Glu Ala Leu His Glu Leu
                                105
Ile Asp Arg Met Asn Val Leu Glu Gln Leu Thr Ala Leu Met Leu Pro
                            120
Trp Gly Cys Asp Val Gln Gly Ser Leu Ser Ser Leu Leu Lys His Leu
                       135
                                            140
Glu Glu Val Pro Gln Leu Val Lys Leu Gly Leu Lys Asn Trp Arg Leu
                   150
                                       155
Thr Asp Thr Glu Ile Arg Ile Leu Gly Ala Phe Phe Gly Lys Asn Pro
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               165
Leu Lys Asn Phe Gln Gln Leu Asn Leu Ala Gly Asn Arg Val Ser Ser
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Asp Gly Trp Leu Ala Phe Met Gly Val Phe Glu Asn Leu Lys
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Thr Tyr Ile Pro Ser Arg Ala Val Ser Leu Phe Phe Asn Trp Lys Gln
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gaa ttc agg act ctg gag gtc aca ctc cgg gat ttc agc aag ttg aat
Glu Phe Arg Thr Leu Glu Val Thr Leu Arg Asp Phe Ser Lys Leu Asn
aag caa gat atc aga tat ctg ggg aaa ata ttc agc tct gcc aca agc
                                                                  144
Lys Gln Asp Ile Arg Tyr Leu Gly Lys Ile Phe Ser Ser Ala Thr Ser
ctc agg ctg caa ata aag aga
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Leu Arg Leu Gln Ile Lys Arg
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<211> 55
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<213> Homo sapiens
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Thr Tyr Ile Pro Ser Arg Ala Val Ser Leu Phe Phe Asn Trp Lys Gln
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Glu Phe Arg Thr Leu Glu Val Thr Leu Arg Asp Phe Ser Lys Leu Asn
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gtc g Val G	gag Hu	ctg Leu	ctg Leu 20	gtc Val	tca Ser	gly ggg	tcc Ser	ctg Leu 25	gaa Glu	ggc Gly	ttc Phe	gag Glu	agt Ser 30	gtc Val	ctg Leu	96
gac t Asp T	gg Trp	ctg Leu 35	ctg Leu	tcc Ser	tgg Trp	gag Glu	gtc Val 40	ctc Leu	tcc Ser	tgg Trp	gag Glu	gac Asp 45	tac Tyr	gag Glu	ggc Gly	144
ttc c	cac His 50	ctc Leu	ctg Leu	ggc Gly	cag Gln	cct Pro 55	ctc Leu	tcc Ser	cac His	ttg Leu	gcc Ala 60	agg Arg	cgc Arg	ctt Leu	ctg Leu	192
gac a Asp T 65	acc Thr	gtc Val	tgg Trp	aat Asn	aag Lys 70	ggt Gly	act Thr	tgg Trp	gcc Ala	tgt Cys 75	cag Gln	aag Lys	ctc Leu	atc Ile	gcg Ala 80	240
gct g Ala A	gcc Ala	caa Gln	gaa Glu	gcc Ala 85	cag Gln	gcc Ala	gac Asp	agc Ser	cag Gln 90	tcc Ser	ccc Pro	aag Lys	ctg Leu	cat His 95	ggc Gly	288
tgc t Cys T	tgg Irp	gac Asp	ccc Pro 100	cac His	tcg Ser	ctc Leu	cac His	cca Pro 105	gcc Ala	cga Arg	gac Asp	ctg Leu	cag Gln 110	agt Ser	cac His	336
cgg ( Arg I	cca Pro	gcc Ala 115	att Ile	gtc Val	agg Arg	agg Arg	ctc Leu 120	cac His	agc Ser	cat His	gtg Val	gag Glu 125	aac Asn	atg Met	ctg Leu	384
gac o Asp l	ctg Leu 130	gca Ala	tgg Trp	gag Glu	cgg Arg	ggt Gly 135	ttc Phe	gtc Val	agc Ser	cag Gln	tat Tyr 140	gaa Glu	tgt Cys	gat Asp	gaa Glu	432
atc a Ile 1 145	agg Arg	ttg Leu	ccg Pro	atc Ile	ttc Phe 150	aca Thr	ccg Pro	tcc Ser	cag Gln	agg Arg 155	gca Ala	aga Arg	agg Arg	ctg Leu	ctt Leu 160	480
gat (	ctt Leu	gcc Ala	acg Thr	gtg Val 165	aaa Lys	gcg Ala	aat Asn	gga Gly	ttg Leu 170	Ala	gcc Ala	ttc Phe	ctt Leu	cta Leu 175	caa Gln	528
cat (	gtt Val	cag Gln	gaa Glu 180	tta Leu	cca Pro	gtc Val	cca Pro	ttg Leu 185	gcc Ala	ctg Leu	cct Pro	ttg Leu	gaa Glu 190	Ala	gcc Ala	576
aca Thr	tgc Cys	aag Lys 195	aag Lys	tat Tyr	atg Met	gcc Ala	aag Lys 200	Leu	agg Arg	acc Thr	acg Thr	gtg Val 205	Ser	gct Ala	cag Gln	624

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														ggc Gly		720
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<b>±</b>				_				_	_	_				tct Ser	_	960
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							_						_	gaa Glu		1104
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														acc Thr		1200
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															ctg Leu		1344
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	_	_		-				_		-				_	gga Gly		1536
	_					_					_		_		aga Arg	-	1584
m Kil kin	Ala	Leu 530	Trp	Gly	Leu	Gly	Met 535	Cys	Cys	Tyr	Val	Phe 540	Ser	Ala	cag Gln	Gln	1632
	Leu 545	Gln	Ala	Ala	Gln	Val 550	Ser	Pro	Asp	Asp	Ile 555	Ser	Leu	Gly	ttc Phe	Leu 560	1680
	Val	Arg	Ala	Lys	Gly 565	Val	Val	Pro	Gly	Ser 570	Thr	Ala	Pro	Leu	gaa Glu 575	Phe	1728
	Leu	His	Ile	Thr 580	Phe	Gln	Cys	Phe	Phe 585	Ala	Ala	Phe	Tyr	Leu 590	gca Ala	Leu	1776
	Ser	Ala	Asp 595	Val	Pro	Pro	Äla	Leu 600	Leu	Arg	His	Leu	Phe 605	Asn	tgt Cys	Gly	1824
	Arg	Pro 610	Gly	Asn	Ser	Pro	Met 615	Ala	Arg	Leu	Leu	Pro 620	Thr	Met	tgc Cys	Ile	1872
	cag	gcc	tcg	gag	gga	aag	gac	agc	agc	gtg	gca	gct	ttg	ctg	cag	aag	1920

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														gcc Ala		2064
			-						_		-	-	_	ggt Gly		2112
														cgg Arg		2160
											_	_	_	cgt Arg 735		2208
											_			ccc Pro		2256
	_	gct	gcc	ctg	gcc	ttt	gtg	cta	cad	cac	ctc	~~~	~~~	000	~+~	2304
	Cys	Ala 755	Ala	Leu	Ala	Phe	Val 760	_	_				~ ~	Pro		2304
	ctg	755 cag	ctg	gac	tac	aac	760 tct	Leu	Gln ggt	His gac	Leu att	Arg 765 ggc	Arg		Val	2352
Ala ctg	ctg Leu 770 ctg	755 cag Gln cct	ctg Leu tgc	gac Asp ctt	tac Tyr	aac Asn 775 gtc	760 tct Ser tgc	Leu gtg Val	Gln ggt Gly gct	His gac Asp	Leu att Ile 780 taa	Arg 765 ggc Gly	Arg gtg Val	Pro gag	Val cag Gln	
ctg Leu 785	ctg Leu 770 ctg Leu	755 cag Gln cct Pro	ctg Leu tgc Cys	gac Asp ctt Leu	tac Tyr ggt Gly 790	aac Asn 775 gtc Val	760 tct Ser tgc Cys	gtg Val aag Lys	Gln ggt Gly gct Ala	gac Asp ctg Leu 795	att Ile 780 taa	Arg 765 ggc Gly ttc Phe T	gtg Val tgg	Pro gag Glu ggc	Cag Gln aac Asn	2352
ctg Leu 785 aga Arg 800	ctg Leu 770 ctg Leu gtg Val	755 cag Gln cct Pro ggt Gly agc	ctg Leu tgc Cys gac Asp	gac Asp ctt Leu gag Glu	tac Tyr ggt Gly 790 ggg Gly 805 tgg	aac Asn 775 gtc Val gcc Ala	tct ser tgc Cys cag Gln	Leu gtg Val aag Lys gcc Ala	Gln ggt Gly gct Ala ctg Leu	gac Asp ctg Leu 795 gct Ala 810	Leu att Ile 780 taa gaa Glu aac	Arg 765 ggc Gly ttc Phe T	gtg Val tgg rp (	gag Glu ggc Sly F	Cag Gln aac asn gat Asp 815	2352

835 . 840 845

gaa gaa ctc tgc ctg gag gag aac cat ctc cag gat gaa ggt gta tgt Glu Glu Leu Cys Leu Glu Glu Asn His Leu Gln Asp Glu Gly Val Cys 850 855 860	2592
tct ctc gca gaa gga ctg aag aaa aat tca agt ttg aaa atc ctg aac Ser Leu Ala Glu Gly Leu Lys Lys Asn Ser Ser Leu Lys Ile Leu Asn 865 870 875	2640
ata aaa att cat gct tcg gga ttc aac aaa ctc ttg gaa agc att ttc Ile Lys Ile His Ala Ser Gly Phe Asn Lys Leu Leu Glu Ser Ile Phe 880 885 890 895	2688
tgc atc ctc ctg gtt gtg gaa gca ttt ttc ctg cag aaa gtt gtc aag Cys Ile Leu Leu Val Val Glu Ala Phe Phe Leu Gln Lys Val Val Lys 900 905 910	2736
att ctt gaa gaa atg gta gtc agt tgg cta gag gtc agg ttg tcc aat  Ile Leu Glu Glu Met Val Val Ser Trp Leu Glu Val Arg Leu Ser Asn  915  920  925	2784
aac tgc atc acc tac cta ggg gca gaa gcc ctc ctg cag gcc ctt gaa Asn Cys Ile Thr Tyr Leu Gly Ala Glu Ala Leu Leu Gln Ala Leu Glu 930 935 940	2832
agg aat gac acc atc ctg gaa gtc tgg ctc cga ggg aac act ttc tct Arg Asn Asp Thr Ile Leu Glu Val Trp Leu Arg Gly Asn Thr Phe Ser	2880
945 950 955	
945 950 955	2928
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cta gag gag gtt gac aag ctc ggc tgc agg gac acc aga ctc ttg ctt Leu Glu Glu Val Asp Lys Leu Gly Cys Arg Asp Thr Arg Leu Leu Leu 960 965 970 975  tgaagtetec gggaggatgt tegteteagt ttgtttgtga geaggetgtg agtttgggcc ccagaggetg ggtgacatgt gttggeagec tetteaaaat gageeetgte etgeetaagg	2988 3048
cta gag gag gtt gac aag ctc ggc tgc agg gac acc aga ctc ttg ctt Leu Glu Glu Val Asp Lys Leu Gly Cys Arg Asp Thr Arg Leu Leu 960 965 970 975  tgaagtetec gggaggatgt tegteteagt ttgtttgtga geaggetgtg agtttgggec ceagaggetg ggtgacatgt gttggeagec tetteaaaat gagecetgte etgeetaagg etgaacttgt tttet  <210> 188 <211> 795 <212> PRT <213> Homo sapiens  <400> 188 Cys Glu Met Cys Ser Gln Glu Ala Phe Gln Ala Gln Arg Ser Gln Leu	2988 3048
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cta gag gag gtt gac aag ctc ggc tgc agg gac acc aga ctc ttg ctt Leu Glu Glu Val Asp Lys Leu Gly Cys Arg Asp Thr Arg Leu Leu Leu 960 965 970 975  tgaagtetee gggaggatgt tegteteagt ttgtttgtga geaggetgtg agtttgggee ccagaggetg ggtgacatgt gttggeagee tetteaaaat gageeetgte etgeetaagg etgaacttgt tttet  <210> 188 <211> 795 <212> PRT <213> Homo sapiens  <400> 188  Cys Glu Met Cys Ser Gln Glu Ala Phe Gln Ala Gln Arg Ser Gln Leu 1 5 10 15  Val Glu Leu Leu Val Ser Gly Ser Leu Glu Gly Phe Glu Ser Val Leu 20 25 30  Asp Trp Leu Leu Ser Trp Glu Val Leu Ser Trp Glu Asp Tyr Glu Gly	2988 3048

65					70					75					80
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Cys	Trp	Asp	Pro 100	His	Ser	Leu	His	Pro 105	Ala	Arg	Asp	Leu	Gln 110	Ser	His
Arg	Pro	Ala 115	Ile	Val	Arg	Arg	Leu 120	His	Ser	His	Val	Glu 125	Asn	Met	Leu
Asp	Leu 130	Ala	Trp	Glu	Arg	Gly 135	Phe	Val	Ser	Gln	Tyr 140	Glu	Cys	Asp	Glu
Ile 145	Arg	Leu	Pro	Ile	Phe 150	Thr	Pro	Ser	Gln	Arg 155	Ala	Arg	Arg	Leu	Leu 160
Asp	Leu	Ala	Thr	Val 165	Lys	Ala	Asn	Gly	Leu 170	Ala	Ala	Phe	Leu	Leu 175	Gln
His	Val	Gln	Glu 180	Leu	Pro	Val	Pro	Leu 185	Ala	Leu	Pro	Leu	Glu 190	Ala	Ala
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Ser	Arg 210	Phe	Leu	Ser	Thr	Tyr 215	Asp	Gly	Ala	Glu	Thr 220	Leu	Cys	Leu	Glu
Asp 225	Ile	Tyr	Thr	Glu	Asn 230	Val	Leu	Glu	Val	Trp 235	Ala	Asp	Val	Gly	Met 240
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305					310		Gln			315					320
				325			Cys	_	330		•			335	
_			340				Asp	345					350		
	_	355		_			Lys 360		_			365	_		
	370					375	Thr				380				
385					390		Lys			395					400
				405			Phe		410	_	_		_	415	
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Lys	Arg	His 435	His	Glu	Pro	Gly	Val 440	Ala	Asp	Arg	Leu	Ile 445	Arg	Leu	Leu
Gln	Glu 450	Thr	Ser	Ala	Leu	His 455	Gly	Leu	Cys	His	Leu 460	Pro	Val	Phe	Ser
Trp 465	Met	Val	Ser	Lys	Cys 470	His	Gln	Glu	Leu	Leu 475	Leu	Gln	Glu	Gly	Gly 480
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Ser Leu Leu Arg Gly Arg Leu Pro Thr Leu Leu His Leu Gly Arg Leu
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Ala Leu Trp Gly Leu Gly Met Cys Cys Tyr Val Phe Ser Ala Gln Gln
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Leu Gln Ala Ala Gln Val Ser Pro Asp Asp Ile Ser Leu Gly Phe Leu
                    550
                                        555
Val Arg Ala Lys Gly Val Val Pro Gly Ser Thr Ala Pro Leu Glu Phe
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Leu His Ile Thr Phe Gln Cys Phe Phe Ala Ala Phe Tyr Leu Ala Leu
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Ser Ala Asp Val Pro Pro Ala Leu Leu Arg His Leu Phe Asn Cys Gly
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Arg Pro Gly Asn Ser Pro Met Ala Arg Leu Leu Pro Thr Met Cys Ile
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Gln Ala Ser Glu Gly Lys Asp Ser Ser Val Ala Ala Leu Leu Gln Lys
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Ala Glu Pro His Asn Leu Gln Ile Thr Ala Ala Phe Leu Ala Gly Leu
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Leu Ser Arg Glu His Trp Gly Leu Leu Ala Glu Cys Gln Thr Ser Glu
                                665
Lys Ala Leu Leu Arg Arg Gln Ala Cys Ala Arg Trp Cys Leu Ala Arg
                            680
Ser Leu Arg Lys His Phe His Ser Ile Pro Pro Ala Ala Pro Gly Glu
                        695
Ala Lys Ser Val His Ala Met Pro Gly Phe Ile Trp Leu Ile Arg Ser
                    710
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Leu Tyr Glu Met Gln Glu Glu Arg Leu Ala Arg Lys Ala Ala Arg Gly
Leu Asn Val Gly His Leu Lys Leu Thr Phe Cys Ser Val Gly Pro Thr
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                                745
Glu Cys Ala Ala Leu Ala Phe Val Leu Gln His Leu Arg Arg Pro Val
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<213> Homo sapiens

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Glu Ser Ile Phe Cys Ile Leu Leu Val Val Glu Ala Phe Phe Leu Gln
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Lys Val Val Lys Ile Leu Glu Glu Met Val Val Ser Trp Leu Glu Val
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Arg Leu Ser Asn Asn Cys Ile Thr Tyr Leu Gly Ala Glu Ala Leu Leu
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Gln Ala Leu Glu Arg Asn Asp Thr Ile Leu Glu Val Trp Leu Arg Gly
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Arg Leu Leu Leu
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ctgctgggaa ggctgctgcc caacctgtgt atccagggct ccagagtcaa gaagggcagc 180
gaagcageee tg etg cag aag get gag eea cae aac etg caa ate aca gea 231
              Leu Gln Lys Ala Glu Pro His Asn Leu Gln Ile Thr Ala
gcc ttc cta gca ggt ctg ttg tcc cag cag cat cgg gac ctg ttg gct
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Ala Phe Leu Ala Gly Leu Leu Ser Gln Gln His Arg Asp Leu Leu Ala
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Ala Cys Gln Val Ser Glu Arg Val Leu Leu Gln Arg Gln Ala Arg Ala
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cgc tcg tgt ctg gcc cac agc ctc cgc gag cac ttc cat tcc atc ccg
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Arg Ser Cys Leu Ala His Ser Leu Arg Glu His Phe His Ser Ile Pro
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cct gcc gtg ccc ggt gag acc aag agc atg cat gct atg ccg ggc ttc
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Pro Ala Val Pro Gly Glu Thr Lys Ser Met His Ala Met Pro Gly Phe
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                                 70
att tgg ctc atc cgt agc ctg tac gag atg cag gag gag cag ttg gcc
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Ile Ser Glu Arg Val Leu Leu Gln Arg Gln Ala Arg Ala Arg Ser Cys

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Pro Gly Glu Thr Lys Ser Met His Ala Met Pro Gly Phe Ile Trp Leu

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